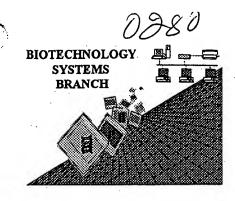
RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following CRF diskette:

Application Serial Number:

09/458,813

Art Unit / Team No.:

OIPE

Date Processed by STIC:

12/30/99

THE ATTACHED PRINTOUT EXPLAINS THE ERRORS DETECTED.

PLEASE BE SURE TO FORWARD THIS INFORMATION TO THE APPLICANTS BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANTS ALONG WITH A NOTICE TO COMPLY or,
- 2) CALLING APPLICANTS AND FAXING THEM A COPY OF THE PRINTOUT WITH A NOTICE TO COMPLY

THIS WILL INSURE THAT THE NEXT SUBMISSION RECEIVED FROM THEM WILL BE ERROR FREE.

IF YOU HAVE ANY FURTHER QUESTIONS, PLEASE CALL:

MARK SPENCER 703-308-4212

Raw Sequence Listing Error Summary

ERROR DETECTED SUGGESTED CORRECTION SERIAL NUMBER: 07/4/38/8/

ATTN	: NEW RULES CASES: F	LEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE
1	Wrapped Nucleics	The number/text at the end of each line "wrapped" down to the next line.
'	· Wappoo Manage	This may occur if your file was retrieved in a word processor after creating it.
		Please adjust your right margin to .3, as this will prevent "wrapping".
^	Managed Aminos	The amino acid number/text at the end of each line "wrapped " down to the next line.
² ——	Wrapped Aminos	This may occur if your file was retrieved in a word processor after creating it.
		Please adjust your right margin to .3, as this will prevent "wrapping".
		Please adjust your right margin to .5, as ans will prover mopping.
3	Incorrect Line Length	The rules require that a line not exceed 72 characters in length. This includes spaces.
4	Misaligned Amino Ácid	The numbering under each 5th amino acid is misaligned. This may be caused by the use of tabs
	Numbering	between the numbering. It is recommended to delete any tabs and use spacing between the numbers.
5	Non-ASCII	This file was not saved in ASCII (DOS) text, as required by the Sequence Rules.
		Please ensure your subsequent submission is saved in ASCII text so that it can be processed.
6	Variable Length	Sequence(s) contain n's or Xaa's which represented more than one residue.
·	- Vallable Lengal	As per the rules, each n or Xaa can only represent a single residue.
		Please present the maximum number of each residue having variable length and
		indicate in the (ix) feature section that some may be missing.
_	Data dia usa 20 "bug"	A "bug" in Patentin version 2.0 has caused the <220>-<223> section to be missing from amino acid
′ —	Patentin ver. 2.0 "bug"	sequence(s) Normally, Patentin would automatically generate this section from the
		previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section
		to the subsequent amino acid sequence.
_		Sequence(s) missing. If intentional, please use the following format for each skipped sequence:
8	Skipped Sequences	
	(OLD RULES)	(2) INFORMATION FOR SEQ ID NO:X: (I) SEQUENCE CHARACTERISTICS:(Do not insert any headings under "SEQUENCE CHARACTERISTICS")
	•	(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X:
		This sequence is intentionally skipped
		Please also adjust the "(iii) NUMBER OF SEQUENCES:" response to include the skipped sequence(s).
9	Skipped Sequences	Sequence(s) missing. If intentional, please use the following format for each skipped sequence.
	(NEW RULES)	<210> sequence id number
		<400> sequence id number
1		000
, J	Use of n's or Xaa's	Use of n's and/or Xaa's have been detected in the Sequence Listing.
	(NEW RULES)	Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
	(,,,,	In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
1	Use of <213>Organism	Sequence(s) are missing this mandatory field or its response.
· —	(NEW RULES)	
2	Use of <220>Feature	Sequence(s) are missing the <220>Feature and associated headings.
	(NEW RULES)	Use of <220> to <223> is MANDATORY if <213>ORGANISM is "Artificial" or "Unknown"
	, :=:: (10220)	Please explain source of genetic material in <220> to <223> section.
		(See "Federal Register," 6/01/98, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of new Rules)
3	Patentia us a a m	Please do not use "Copy to Disk" function of Patentin version 2.0. This causes a corrupted
-	Patentin ver. 2.0 "bug"	file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing).
		Instead, please use "File Manager" or any other means to copy file to floppy disk.
	,	AKS-Biotechnology Systems Branch- 5/15/99

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<220> FEATURE:

RAW SEQUENCE LISTING PATENT APPLICATION US/09/458,813

DATE: 12/30/1999 TIME: 14:42:54

Input Set: I458813.RAW

This Raw Listing contains the General Information Section and up to first 5 pages.

Corrected Diskette Needer Does Not Comply 1 <110> APPLICANT: Murphy, Brian R. 2 Collins, Peter L. 3 Durbin, Anna P. 4 Skiadopoulos, Mario H. 5 Tao, Tao <120> TITLE OF INVENTION: USE OF RECOMBINANT LIVE-ATTENUATED PARAINFLUENZA VIRUS 6 7 (PIV) AS A VECTOR TO PROTECT AGAINST DISEASE CAUSED BY 8 PIV AND RESPIRATORY SYNCYTIAL VIRUS (RSV) 9 <130> FILE REFERENCE: 17634-000330 10 <140> CURRENT APPLICATION NUMBER: US/09/458,813 11 <141> CURRENT FILING DATE: 1999-12-10 <150> EARLIER APPLICATION NUMBER: 09/083,793 12 13 <151> EARLIER FILING DATE: 1998-05-22 <150> EARLIER APPLICATION NUMBER: 60/047,575 <151> EARLIER FILING DATE: 1997-05-23 <150> EARLIER APPLICATION NUMBER: 60/059,385 <151> EARLIER FILING DATE: 1997-09-19 <160> NUMBER OF SEQ ID NOS: 30 18 <170> SOFTWARE: PatentIn Ver. 2.1 20 <210> SEQ ID NO 1 21 <211> LENGTH: 42 <212> TYPE: DNA 22 23 <213> ORGANISM: Artificial Sequence 24 <220> FEATURE: 25 <223> OTHER INFORMATION: Description of Artificial Sequence: Nucleotide 26 insert to conform inserted sequence to rule of 27 six. 28 <400> SEQUENCE: 1 29 cttaagaata tacaaataag aaaaacttag gattaaagag cg 42 30 <210> SEQ ID NO 2 <211> LENGTH: 36 31 <212> TYPE: DNA 33 <213> ORGANISM: Artificial Sequence <220> FEATURE: <223> OTHER INFORMATION: Description of Artificial Sequence: Nucleotide insert to conform inserted sequence to rule of 37 six. 38 <400> SEQUENCE: 2 39 gatccaacaa agaaacgaca ccgaacaaac cttaag 36 40 <210> SEQ ID NO 3 41 <211> LENGTH: 101 42 <212> TYPE: DNA 43 <213> ORGANISM: Artificial Sequence

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        46
        47
                   six.
        48
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        49
                   tataattgag gaaggaccca atagacaaat ccaaattcga g
                                                                                       101
        50
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             <211> LENGTH: 79
        52
        53
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             <213> ORGANISM: Artificial Sequence
        54
        55
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        56
                   insert to conform inserted sequence to rule of
        57
        58
                   six.
        59
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        61
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        63
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        66
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        67
                   insert to conform inserted sequence to rule of
        68
        69
             <400> SEQUENCE: 5 see Len 10 on Eva Summary Sheet
        70
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        71
W-->
        72
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        73
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        74
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        77
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        79
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        80
                   six.
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        81
                                                                                       44
        82
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        88
        89
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        90
                   six.
             <400> SEQUENCE: 7
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                                                                                       6
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        93
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PATENT APPLICATION US/09/458,813

Input Set: I458813.RAW

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 95
 96
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 97
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 98
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 99
100
            six.
101
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102
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103
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110
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112
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                                                                                6
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119
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120
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121
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122
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130
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132
133
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137
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138
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139
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140
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                                                      > den 10
141
            six.
142
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                                                                                34
143
           agacaatntc ntndrmrvsa gtntcntndn ntag
144
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Input Set: 1458813.RAW

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174		gtccttcc	68
175		SEQ ID NO 16	
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177		TYPE: DNA	
178		ORGANISM: Artificial Sequence	
179		FEATURE:	
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182	-400-	SIX.	
183	<400>	SEQUENCE: 16	60
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186	-210-	gtccttcc SEQ ID NO 17	00
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188		TYPE: DNA	
189		ORGANISM: Artificial Sequence	
190		FEATURE:	
191		OTHER INFORMATION: Description of Artificial Sequence: Nucleotide	
192	~2237	insert to conform inserted sequence to rule of	
193		six.	
194	<400>	SEQUENCE: 17	

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TIME: 14:42:54

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RAW SEQUENCE LISTING
PATENT APPLICATION US/09/458,813

Input Set: I458813.RAW

28 195 ccatgtaatt gaatccccca acactagc 196 <210> SEQ ID NO 18 197 <211> LENGTH: 28 <212> TYPE: DNA 198 <213> ORGANISM: Artificial Sequence 199 200 <220> FEATURE: <223> OTHER INFORMATION: Description of Artificial Sequence: Nucleotide 201 202 insert to conform inserted sequence to rule of 203 six. 204 <400> SEQUENCE: 18 28 205 cggataaacg cgttctacaa agataacc 206 <210> SEQ ID NO 19 207 <211> LENGTH: 28 208 <212> TYPE: DNA 209 <213> ORGANISM: Artificial Sequence 210 <220> FEATURE: 211 <223> OTHER INFORMATION: Description of Artificial Sequence: Nucleotide insert to conform inserted sequence to rule of 212 213 six. 214 <400> SEQUENCE: 19 28 215 cggataaacg cgttctacaa agataacc 216 <210> SEQ ID NO 20 <211> LENGTH: 23 217 <212> TYPE: DNA 218 219 <213> ORGANISM: Artificial Sequence 220 <220> FEATURE: <223> OTHER INFORMATION: Description of Artificial Sequence: Nucleotide 221 222 insert to conform inserted sequence to rule of 223 six. 224 <400> SEQUENCE: 20 gggccatgga agattacagc aat 225 23 226 <210> SEQ ID NO 21 227 <211> LENGTH: 25 <212> TYPE: DNA 228 229 <213> ORGANISM: Artificial Sequence 230 <220> FEATURE: <223> OTHER INFORMATION: Description of Artificial Sequence: Nucleotide 231 232 insert to conform inserted sequence to rule of 233 six. 234 <400> SEQUENCE: 21 235 25 caataagctt aaagcattag ttccc 236 <210> SEQ ID NO 22 237 <211> LENGTH: 31 238 <212> TYPE: DNA 239 <213> ORGANISM: Artificial Sequence 240 <220> FEATURE: 241 <223> OTHER INFORMATION: Description of Artificial Sequence: Nucleotide 242 insert to conform inserted sequence to rule of 243 six. 244 <400> SEOUENCE: 22

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VERIFICATION SUMMARY
PATENT APPLICATION US/09/458,813

DATE: 12/30/1999 TIME: 14:42:54

Input Set: I458813.RAW

Line	? Error/Warning				Original Text						
							required required	nnnaggaaaa ctccctnnn	gggaaatata	aaaacttagg	agtaaagt
82	W	"N"	or	"Xaa"	used:	Feature	required required	nnnaggaaaa	gggaacgcgt ntndrmrvsa	-	